

Notice of Allowability	Application No.	Applicant(s)	
	10/614,954	HOHN ET AL.	
	Examiner Medina A. Ibrahim	Art Unit 1638	

-- **The MAILING DATE of this communication appears on the cover sheet with the correspondence address--**

All claims being allowable, PROSECUTION ON THE MERITS IS (OR REMAINS) CLOSED in this application. If not included herewith (or previously mailed), a Notice of Allowance (PTOL-85) or other appropriate communication will be mailed in due course. **THIS NOTICE OF ALLOWABILITY IS NOT A GRANT OF PATENT RIGHTS.** This application is subject to withdrawal from issue at the initiative of the Office or upon petition by the applicant. See 37 CFR 1.313 and MPEP 1308.

1. This communication is responsive to the amendment of 05/09/05.
2. The allowed claim(s) is/are 1-5.
3. The drawings filed on 07 July 2003 are accepted by the Examiner.
4. Acknowledgment is made of a claim for foreign priority under 35 U.S.C. § 119(a)-(d) or (f).
 - a) All
 - b) Some*
 - c) None
 of the:
 1. Certified copies of the priority documents have been received.
 2. Certified copies of the priority documents have been received in Application No. _____.
 3. Copies of the certified copies of the priority documents have been received in this national stage application from the International Bureau (PCT Rule 17.2(a)).

* Certified copies not received: _____.

Applicant has THREE MONTHS FROM THE "MAILING DATE" of this communication to file a reply complying with the requirements noted below. Failure to timely comply will result in ABANDONMENT of this application.
THIS THREE-MONTH PERIOD IS NOT EXTENDABLE.

5. A SUBSTITUTE OATH OR DECLARATION must be submitted. Note the attached EXAMINER'S AMENDMENT or NOTICE OF INFORMAL PATENT APPLICATION (PTO-152) which gives reason(s) why the oath or declaration is deficient.
6. CORRECTED DRAWINGS (as "replacement sheets") must be submitted.
 - (a) including changes required by the Notice of Draftsperson's Patent Drawing Review (PTO-948) attached
 - 1) hereto or 2) to Paper No./Mail Date _____.
 - (b) including changes required by the attached Examiner's Amendment / Comment or in the Office action of Paper No./Mail Date _____.

Identifying indicia such as the application number (see 37 CFR 1.84(c)) should be written on the drawings in the front (not the back) of each sheet. Replacement sheet(s) should be labeled as such in the header according to 37 CFR 1.121(d).
7. DEPOSIT OF and/or INFORMATION about the deposit of BIOLOGICAL MATERIAL must be submitted. Note the attached Examiner's comment regarding REQUIREMENT FOR THE DEPOSIT OF BIOLOGICAL MATERIAL.

Attachment(s)

1. Notice of References Cited (PTO-892)
2. Notice of Draftsperson's Patent Drawing Review (PTO-948)
3. Information Disclosure Statements (PTO-1449 or PTO/SB/08),
Paper No./Mail Date _____
4. Examiner's Comment Regarding Requirement for Deposit
of Biological Material
5. Notice of Informal Patent Application (PTO-152)
6. Interview Summary (PTO-413),
Paper No./Mail Date _____.
7. Examiner's Amendment/Comment
8. Examiner's Statement of Reasons for Allowance
9. Other _____.

EXAMINER'S AMENDMENT

An examiner's amendment to the record appears below. Should the changes and/or additions be unacceptable to applicant, an amendment may be filed as provided by 37 CFR 1.312. To ensure consideration of such an amendment, it MUST be submitted no later than the payment of the issue fee.

Authorization for this examiner's amendment was given in a telephone interview with Gregory Warren on 06/25/05.

The application has been amended as follows:

In The Title:

The title has been replaced with the following:

--- ISOLATED NUCLEIC ACIDS ENCODING TRICHOTHECENE 3-O-ACETYLTRANSFERASES ----.

In The Abstract:

The abstract has been replaced with the following:

---- The present invention discloses isolated polynucleotides encoding polypeptides having trichothecene 3-O-acetyltransferase activity, recombinant vectors and host cell comprising said polynucleotides----.

In The claims:

In claims 3-5, "gene" has been replaced with ---DNA----.

Reasons for allowance

The following is an examiner's statement of reasons for allowance: the prior art does not teach or reasonably suggest a nucleotide sequence that is 100% identical to

SEQ ID NO: 5 or a nucleotide sequence encoding SEQ ID NO: 6. The closest prior art is Kimura et al (J. Biol. Chem. (1998), 273 (3): 1654-1661, Applicant's IDS) who teach a nucleotide sequence having 99% sequence identity with Applicant's SEQ ID NO: 5. See attached sequence alignment results (Exhibit A/B) between SEQ ID NO: 5/6 and the prior art sequences.

Any comments considered necessary by applicant must be submitted no later than the payment of the issue fee and, to avoid processing delays, should preferably accompany the issue fee. Such submissions should be clearly labeled "Comments on Statement of Reasons for Allowance."

Contact information

Any inquiry concerning this communication or earlier communications from the examiner should be directed to Medina A. Ibrahim whose telephone number is (571) 272-0797. The Examiner can normally be reached Monday -Thursday from 8:00AM to 5:30PM and every other Friday from 9:00AM to 5:00 PM. Before and after final responses should be directed to fax nos. (703) 872-9306 and (703) 872-9307, respectively.

If attempts to reach the Examiner by telephone are unsuccessful, the Examiner's supervisor, Dr. Amy Nelson, can be reached at (571) 272-0804.

Information regarding the status of an application may be obtained from the Patent Application Information Retrieval (PAIR) system. Status information for published applications may be obtained from either Private PAIR or Public PAIR. Status information for unpublished applications is available through Private PAIR only. For more information about the PAIR system, see <http://pair-direct.uspto.gov>. Should you have questions on access to the Private PAIR system, contact the Electronic Business Center (EBC) at 866-217-9197 (toll-free).

For all other customer support, please call the USPTO Call Center (UCC) at 800-786-9199.

6/25/05
Mai

MEDINA A. IBRAHIM
PATENT EXAMINER
Medina A. Ibrahim
1638

06 May 2005

Alignment Results

Alignment: Global DNA alignment against reference molecule
 Parameters: Scoring matrix: Linear (Mismatch 2, OpenGap 4, ExtGap 1)

Reference molecule: SEQIDNO:5, Region 1-1356

Number of sequences to align: 2

Settings: Similarity significance value cutoff: >= 60%



Summary of Percent Matches:

Reference:	SEQIDNO:5	1 -	1356	(1356 bps)	--
Sequence 2:	Kimura Tri101	135 -	1490	(1356 bps)	99%

SEQIDNO:5 Kimura Tri10	1 135	atggcttcaagatacagctcgacaccctcgccagctaccaggcctcttcgatctac atggcttcaagatacagctcgacaccctcgccagctaccaggcctcttcgatctac
SEQIDNO:5 Kimura Tri10	61 195	acccaaatcagtctccttacccgtctgtatccctcaatatcccactattgtcagc acccaaatcagtctccttacccgtctgtatccctcaatatcccactattgtcagc
SEQIDNO:5 Kimura Tri10	121 255	accttcgagcaaggcttaagcgcttccgaagccgtccatgggtcgccaggcaggc accttcgagcaaggcttaagcgcttccgaagccgtccatgggtcgccaggcaggc
SEQIDNO:5 Kimura Tri10	181 315	aaagccgagggcattagcgagggaaacacaggaacttccttatcgccctttgaggac aaagccgagggcattagcgagggaaacacggaaacttccttatcgccctttgaggac
SEQIDNO:5 Kimura Tri10	241 375	gttcctcgttgttagtggaaacactccgcgtatccctcagccccacgatcgagggt gttcctcgttgttagtggaaacactccgcgtatccctcagccccacgatcgagggt
SEQIDNO:5 Kimura Tri10	301 435	atgagaaaggcggataccctatggcgatgtttgacgagaacatcatcgcccaaggaa atgagaaaggcggataccctatggcgatgtttgacgagaacatcatcgcccaaggaa
SEQIDNO:5 Kimura Tri10	361 495	acgttacctattggacctggtaactggccgacgacccaaagcctgtatttattgcag acgttacctattggacctggtaactggccgacgacccaaagcctgtatttattgcag
SEQIDNO:5 Kimura Tri10	421 555	ctcaacttcatcaaggcggactcatccctactgtcaacggacacgacgggtctatggat ctcaacttcatcaaggcggactcatccctactgtcaacggacacgacgggtctatggat
SEQIDNO:5 Kimura Tri10	481 615	atggtaggccaagatgcggtagtccgtctactctccaaaggcgtccgtaacgaccatc atggtaggccaagatgcggtagtccgtctactctccaaaggcgtccgtaacgaccatc
SEQIDNO:5 Kimura Tri10	541 675	accgaagagggaaatgacggccatgaacctcgatcgcaagacgatagttccttacctt accgaagagggaaatgacggccatgaacctcgatcgcaagacgatagttccttacctt
SEQIDNO:5 Kimura Tri10	601 735	aactatacgattggcccgaggttagatcatcgatgtcaaaactgtatgttagctgg aactacacgattggcccgaggttagatcatcgatgtcaaaactgtatgttagctgg
SEQIDNO:5 Kimura Tri10	661 795	gacgctgttctcacgcccgtcaagtgcacgtggcggtttcacattcagccccaaagg gacgctgttctcacgcccgtcaagtgcacgtggcggtttcaattcagccccaaagg
SEQIDNO:5 Kimura Tri10	721 855	atgtcagagctcaaggatgtctaccaggactcttgcacgcataacaaagtgcgt atgtcagagctcaaggatgtctaccaggactcttgcacgcataacaaagtgcgt
SEQIDNO:5 Kimura Tri10	781 915	actgacgatgtctttcgccgtcatctggaaatcgcccttcgcgtgcgtctcgaaa actgacgatgtctttcgccgtcatctggaaatcgcccttcgcgtgcgtctcgaaa
SEQIDNO:5 Kimura Tri10	841 975	atcgatggctctgcacccatccggagttctggcggtgttgcgtgcaccggcaatgg atcgatggctctgcacccatccggagttctggcggtgttgcgtgcaccggcaatgg
SEQIDNO:5 Kimura Tri10	901 1035	gtctcgaacaactaccaggccttcaaaaacatgacctaccacaactcgaccatcg gtctcgaacaactaccaggccttcaaaaacatgacctaccacaactcgaccatcg
SEQIDNO:5 Kimura Tri10	961 1095	gaaatcgccaaacgaggtaactcgccgcaacacgcatcagccttcgtt gaaatcgccaaacgaggtaactcgccgcaacacgcatcagccttcgtt

SEQIDNO:5 1021 gcgagcatgcgccagcgaacaagaggctcgcgacgtacctgcacaacaaccccgacaag
Kimura Tri10 1155 gcgagcatgcgccagcgaacaagaggctcgcgacgtacctgcacaacaaccccgacaag

SEQIDNO:5 1081 tccaacgttatccctgacggctgatcgacggaccatctaccagcgtcatgctgagttttgg
Kimura Tri10 1215 tccaacgttatctcgacggctgatcgacggaccatctaccagcgtcatgctgagttttgg

SEQIDNO:5 1141 gccaagggtggactctgggattacgactttggctcgactggtaagcccgagactgtg
Kimura Tri10 1275 gccaagggtcgactctgggattacgactttggctcgactggtaagcccgagactgtg

SEQIDNO:5 1201 agacggccaatcttgagcctgtgagagcttgcgtactttatgcccagaaggcctgat
Kimura Tri10 1335 agacggccaatcttgagcctgtgagagcttgcgtactttatgcccagaaggcctgat

SEQIDNO:5 1261 ggcgaggttctgtgcggcgcttctcgagggatgaggatatggaccgattgaaggcggat
Kimura Tri10 1395 ggcgaggttctgtgcggcgcttctcgagggatgaggatatggaccgattgaaggcggat

SEQIDNO:5 1321 aaggagtggaccaagtatgcgcagtacgttggtag
Kimura Tri10 1455 aaggagtggaccaagtatgcgcagtacgttggtag

06 May 2005

Alignment Results

Exhibit B

Alignment: Global Protein alignment against reference molecule
Parameters: Scoring matrix: BLOSUM 62

Reference molecule: SEQIDNO:6, Region 1-451

Number of sequences to align: 2

Settings: Similarity significance value cutoff: >= 60%



Summary of Percent Matches:

Reference:	SEQIDNO:6	1 - 451 (451 aa)	--
Sequence 2:	Kimura Tri101 Pro	1 - 451 (451 aa)	99%

SEQIDNO:6	1 mafkiqltdtigqlpgllsiytqisllypvsds	sqyptivstfeqglkrfseavpwwagqv
Kimura Tri10	1 mafkiqltdtigqlpgllsiytqisllypvsdp	sqyptivstfeqglkrfseavpwwagqv
SEQIDNO:6	61 kaegiseqntgtsfivp fedvprvvvkdlrddps	aptiegmrkagypmamfdeniiaprk
Kimura Tri10	61 kaegiseqntgtsfivp fedvprvvvkdlrddps	aptiegmrkagypmamfdeniiaprk
SEQIDNO:6	121 tlpigpgtgpddpkpvillqlnfikggli	tvngqhgamdmvgqdadvirllskacrndpf
Kimura Tri10	121 tlpigpgtgpddpkpvillqlnfikggli	tvngqhgamdmvgqdadvirllskacrndpf
SEQIDNO:6	181 teeemtamnlrktivpylenytigpevdhqiv	kadvaggdavltltpvsaswafftfspka
Kimura Tri10	181 teeemtamnlrktivpylenytigpevdhqiv	kadvaggdavltltpvsaswaffkifspka
SEQIDNO:6	241 mselkdaatktidastkf vst ddalsafiwks	sasrvrleridgsaptefcravdarpamg
Kimura Tri10	241 mselkdaatktldastkf vst ddalsafiwks	sasrvrleridgsaptefcravdarpamg
SEQIDNO:6	301 vsnnypgllqnmtynstigeianeslgatasrl	rseldpasmrqrtrglatylhnnpdk
Kimura Tri10	301 vsnnypgllqnmtynstigeianeslgatasrl	rseldpasmrqrtrglatylhnnpdk
SEQIDNO:6	361 snvsltadadpstsvmlsswakvglwdydfg	lgkpetvrrpifepveslmyfmpkkpd
Kimura Tri10	361 snvsltadadpstsvmlsswakvglwdydfg	lgkpetvrrpifepveslmyfmpkkpd
SEQIDNO:6	421 gefcaalslrdedmdrlkadkewtkyaqyvg	
Kimura Tri10	421 gefcaalslrdedmdrlkadkewtkyaqyvg	